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IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

In re Application of: Lal *et al.*

Art Unit: 1645

Application No. 09/763,397

CERTIFICATE OF MAILING

Filed: February 16, 2001


For: RECOMBINANT MULTIVALENT MALARIAL
VACCINE AGAINST PLASMODIUM
FALCIPARUM

Examiner: Vanessa L. Ford

Date: June 30, 2004

I hereby certify that this paper and the documents referred to as being attached or enclosed herewith are being deposited with the United States Postal Service as First Class Mail in an envelope addressed to: COMMISSIONER FOR PATENTS, P.O. BOX 1450, ALEXANDRIA, VA 22313-1450 on the date shown below.

Attorney
for Applicant(s)



Date Mailed June 30, 2004

COMMISSIONER FOR PATENTS
P.O. BOX 1450
ALEXANDRIA, VA 22313-1450

DECLARATION UNDER 37 C.F.R. § 1.131

We, Ya Ping Shi, Altaf A. Lal and Seyed E. Hasnain hereby declare as follows:

1. We are the co-inventors of the subject matter described and claimed by the patent application referenced above, *i.e.*, United States application No. 09/763,397 (hereafter the '397 application). We were employed by the Centers for Disease Control and Prevention (CDC), the assignee of the '397 application, which is located in Atlanta, Georgia, while developing the invention described and claimed in the referenced application.

2. We understand that claims pending in the present application have been rejected in view of Gilbert *et al.*, *Nature Biotechnology*, 15: 1280-1284, 1997. We understand that Gilbert *et al.*, has been cited as allegedly anticipating certain claims pending in the referenced application, or, in the alternative, as allegedly rendering the claimed embodiments obvious.

3. The publication date of Gilbert *et al.*, is November 1997. United States Provisional Application No. 60/097,703 was filed on August 21, 1998. However, we invented the subject matter covered by the claims pending in the '397 application well prior to the November 1997 date that Gilbert *et al.*, became available as a reference.

4. Accompanying this Declaration as Exhibit A are photocopies of pages from Dr. Shi's laboratory research notebook. These copies are true and accurate facsimile copies of

photocopies of the corresponding pages from Dr. Shi's laboratory notebooks. All dates stated on these pages have been redacted.

5. All entries on the notebook pages of Exhibit A were made prior to November 1997.

6. The ideas and concepts demonstrated by Exhibit A arose from work conducted for the CDC in Atlanta, GA. These ideas and concepts are embodied in the claims of the '397 application. Thus, conception and reduction to practice of the invention recited in the claims of the '397 application, as discussed in more detail below, occurred in the United States of America prior to November 1997.

7. Exhibit A consists of 21 pages of laboratory notebook pages. Exhibit B consists of one page of CDC Biotechnology Core Facility Records. The contents of these pages of Exhibits A and B, and pertinent statements made on these pages are discussed below.

A. Exhibit B is a record from the CDC Biotechnology Core Facility showing the dates of a request for oligonucleotide synthesis, and the sequences of the requested oligonucleotides. These requests were made prior to November 1997. These oligonucleotides were used as is depicted in Figure 2 of the specification to amplify the synthetic vaccine antigen gene construct using Polymerase Chain Reactions (PCRs). The oligonucleotides of Exhibit B consist of both forward and reverse complementary sequences of SEQ ID NO: 1 of the application, with overlapping sequences acting as primers for the amplification in either the forward or reverse direction.

B. Pages 1-6 of Exhibit A display the planning strategy for the PCR synthesis of the synthetic gene construct. Set forth are relevant calculations for PCR reactions and primers used to generate quantities of the synthetic gene construct. Also shown are electrophoresis gels used to visually confirm the size of PCR-generated products.

1) Page 1 shows the calculation and strategy for serial PCRs. As is set forth at the top of page one, "AA" was short hand for the PCR reaction involving oligonucleotides G0, GL, G1, and G2 of Exhibit B. "BB" was short hand for the PCR reaction involving oligonucleotides G3-G6, and "CC" was short hand for the PCR reaction involving oligonucleotides G7-G12.

2) Pages 2-5 show several experiments, ending with success as indicated by the comment “works well” on page 5. Reactions DD-II as depicted were successive rounds of PCR that joined the amplified fragments into the final synthetic gene construct.

C. Page 6 shows an electrophoresis gel of four samples at different concentrations from PCR reactions (the central four bands of the gel). Next to the gel is the comment “good!” indicating that the size of the band corresponding to the PCR product in each sample appeared to be the correct size.

D. Pages 7-10 shows that the PCR product was isolated and purified from the electrophoresis gel shown on page 6. Next, the purified product was cut with restriction endonucleases with BamH1 and Not1 (shown as steps #3 and #4 in Figure 2) for cloning into the expression vector pBluescript. The resulting sequence was SEQ ID NO: 1 of the application. Also shown on page 8 are the ligation reaction conditions for the ligation reaction, followed by restriction endonuclease reactions to evaluate the success of the ligation. The vector containing the fragment was then transformed into cells and plated onto agar plates. Positive clones were identified by their white color, indicating that the blue color-producing gene characteristic of a vector without a cloned segment had been interrupted with a cloned fragment. Page 10 sets forth the conditions for the PCR reaction to confirm that the correct gene fragment had been ligated into the vector (*i.e.*, to identify positive clones). The notations indicates that seventeen positive clones (numbers 1-4, 6, 8, 17, 21, 22, 25-27, 31, 33, 36, 39 and 40) were identified.

E. Page 11 shows an electrophoresis gel displaying samples of the PCR products.

F. Page 12 shows a single and double digesting experiment to confirm that the cloned fragment was properly oriented and was of the correct size. Clones 3, 26, and 33 were discarded by this experiment, leaving fourteen correct clones.

G. Page 13 shows the methods for the transformation of two plasmids, pBacPAK8 and pBacPAK9 with the synthetic gene construct for expression in Baculovirus. Also shown is an electrophoresis gel displaying samples of digested and undigested plasmid.

H. Page 14 shows another electrophoresis gel containing samples of DNA that were purified and digested with restriction endonucleases Not1 and BamH1, to confirm that the cloning into the Baculovirus expression vectors had been successful. The notation “orders are no problem” indicates that the clones were correctly oriented, and the statement “confirm 11, 20, 63 clones are true clones” indicates that these clones were considered to be successful. The

depicted gel shows the results of restriction endonuclease digestion showing the two bands of each clone (lanes 4, 5 and 8, respectively). Clone number 20 was identified as the clone that would be sequenced to confirm correctness at the molecular level. The sequencing indicated that clone 20 contained a single mutation. Therefore, a second clone, number 63 was sequenced. The sequencing indicated that clone 63 also contained a single mutation. In comparing the location of the mutations in clones 20 and 63, the mutations were found to be located in different segments. Thus, subsequent experiments was performed to generate a subclone that would contain the correct segments of clones 20 and 63.

I. Page 15 sets forth conditions for methylation experiments, which were run to protect restriction endonuclease sites in the vector. Following these reactions, the correct segments of clones 20 and 63 (as shown on the bottom of pg. 15) were excised and ligated into the Baculovirus expression plasmid pBacPAK8.

J. Page 16 shows a basic diagram of the recombinant vaccine antigen gene as cloned into the Baculovirus expression vectors. As shown, the construct contains portions of both clones 20 and 63.

K. Page 17 shows a gel wherein the products of the second ligation reaction of the correct segments of constructs 20 and 63 into the pBacPAK8 expression vector were run to confirm the size of the construct.

L. Page 18 shows the success of a ligation experiment, as confirmed by visualization of the bands on the electrophoresis gel. The statement that “clones 21, 31, 33, 35 are positive” indicates that the ligation reaction was successful.

M. Page 19 shows an electrophoresis gel displaying the results of BamH1 restriction endonuclease digestion, to confirm the successful clones.

N. Page 20 shows an electrophoresis gel displaying the results of BamH1 and Not1 restriction endonuclease digestion. The notation “save clone 31A and 31B” indicates that these clones were successful. These results were subsequently confirmed by sequencing clones 31A and 31B.

O. Page 21 contains the notation “miniprep for sending product to Hassian (sic).” This refers to co-inventor Dr. Seyed Hasnain, who tested the expression of the synthetic gene construct in the Baculovirus expression system.

8. All statements made herein and of our own knowledge are true and all statements made on information are believed to be true. Furthermore, these statements were made with the knowledge that willful false statements and the like are punishable by fine or imprisonment, or both, under Section 1001 of Title 18 of the United States Code, and that any such willful false statements made may jeopardize the validity of the application or any patent issuing thereon.

06/14/2004
Date

Ya Ping Shi
Name Ya Ping Shi

Date

Name Altaf A. Lal

Date

Name Seyed E. Hasnain

8. All statements made herein and of our own knowledge are true and all statements made on information are believed to be true. Furthermore, these statements were made with the knowledge that willful false statements and the like are punishable by fine or imprisonment, or both, under Section 1001 of Title 18 of the United States Code, and that any such willful false statements made may jeopardize the validity of the application or any patent issuing thereon.

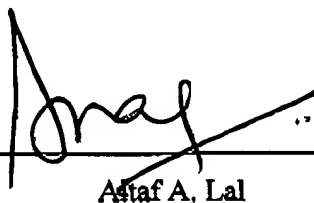
Date

June 14th, 2004

Date

Name

Ya Ping Shi



Name

Asaf A. Lal

Date

Name

Sayed E. Hasnain

8. All statements made herein and of our own knowledge are true and all statements made on information are believed to be true. Furthermore, these statements were made with the knowledge that willful false statements and the like are punishable by fine or imprisonment, or both, under Section 1001 of Title 18 of the United States Code, and that any such willful false statements made may jeopardize the validity of the application or any patent issuing thereon.

Date

Name Ya Ping Shi

Date

Name Altaf A. Lal

June 21, 2007
Date

Sayed E. Hasnain
Name Seyed E. Hasnain

EXHIBIT A

First PCR

AA: G70 - G72
 BB: G73 - G76
 CC: G77 - G712

Tm
 52.5
 50-56
 60-66

94°C 5min
 94°C 45"
 45°C 1min
 72°C 1.5min

8 cycle (p139)

500ng/cath.

AA: 2x4 = 81.4 65.5

BB: 2x4 = 81.4 65.5

CC: 2x6 = 121.4 61.5

16ul dNTP
 10ul Buffer
 0.5ul Taq
 26.5ul

Second PCR

53.5

16ul dNTP
 10ul Buffer
 0.5ul Taq

5ul orig
 5ul 0.182
 46.5

94°C 5min.

94°C 45"

45°C 1min

72°C 1.5min

25 cycle
 (P141)

AA: DD1 1ul 52.5+5+5
 DD2 2.5ul 51+5+5 → G0
 DD3 5ul 48.5+5+5 → G2
 DD4 10ul 43.5+5+5
 BB: EE1 1ul 52.5+5
 EE2 2.5ul 51+5 → G3
 EE3 5ul 48.5+5 → G6
 EE4 10ul 43.5+5

C: FF1 1ul 52.5+5
 FF2 2.5ul 51+5 → G7
 FF3 5ul 48.5+5 → G12
 FF4 10ul 43.5+5

Redo CC₀: G7 - G12 = 12ul

dNTP	16ul	c.; Taq
10x Buffer	10ul	
H ₂ O	61.5ul	
	<u>10ul</u>	

94°C 5min
94°C 45"
40°C 1min
72°C 2min

} 8 cycle.

CC₁: G7 - G8 (only do second PCR) = 4ul + 69.5ul H₂O
CC₂: G9 - G12 2x4 = 8ul + H₂O 65.5

Do SOE: G6 - G6

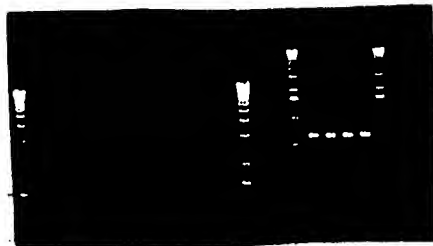
	DD ₁ + EE ₁	H ₂ O	16ul dNTP
G ₁ G ₁	1ul + 1ul = 2ul	61.5	10ul Buffer
G ₂ G ₂	2.5ul + 2.5ul = 5ul	58.5	5ul G ₀
G ₃ G ₃	5ul + 5ul = 10ul	53.5	5ul G ₆
G ₄ G ₄	10ul + 10ul = 20ul	43.4	0.5 Taq
			<u>36.5</u>

program 141

FF ₁	} CC ₁	primers	67.5	H ₂ O	16ul dNTP
FF ₂		G7	1ul	62.5	10ul Buffer
FF ₃		G12	2.5ul	61	oligos 10ul
FF ₄			5ul	58.5	Taq 0.5
FF ₅	} CC ₂	primers	1ul		94°C 5min
FF ₆		G9	2.5ul		94°C 45"
FF ₇		G12	5ul		40°C 1min
FF ₈			10ul		72°C 2min
					<u>36.5</u>

cycles #41

Result: G₅ - 4



IF-3 did not work probably because oligo ?

Prepare new temp oligo G₉ - G₁₂ also AG105.

Redo: $CC'_2 \rightarrow CC''_2$ and CC''_3 .

23.5

CC''_2 G9 G10 G11 G12 $\times 2 = 8 \mu l$. 65.5

CC''_3 G9 G10 G11 $Al1064 \times 2 = 8 \mu l$. 65.5

\times
works well

16 μl dNTP.

10 μl Buffer.

0.5 Tag

Same to before.

Second PCR.

FF''₁
FF''₂
FF''₃
FF''₄

} CC''_2

primers

G9

G12

1 μl

2.5 μl

5 μl

10 μl

H₂O

62.5

61

58.5

53.5

16 μl dNTP

10 μl buffer

0.1 μl Taq

0.5 μl Tag

FF''₅
FF''₆
FF''₇
FF''₈

} CC''_3

G9

Al1064

works well

Same to before.

1141





114



11



SoE for G17 - G11 + AL1065

	CC ₁	FF ₅	H ₂ O	65.5	16ul dNTP
HH ₁	1ul	+ 1ul	61.5		10ul buffer
HH ₂	2.5ul	+ 2.5ul	58.5		5ul G7
HH ₃	5ul	+ 5ul	53.5		5ul AL1065
HH ₄	10ul	+ 10ul	43.5		0.5 T99

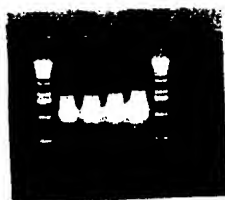
36.5

program #41

	G171	+ HH ₁	H ₂ O	63.5	16ul dNTP
II ₁	1ul 1ul	+ 1ul	61.5		10ul buffer
II ₂	2.5ul	+ 2.5ul	58.5		5ul AL1064
II ₃	5ul	+ 5ul	53.5		5ul AL1065
II ₄	10ul	+ 10ul	43.5		0.5 T99

36.5

program #41



good!

Further cleaning and cloning,
sequencing.

A: Run gel and cut ~~and~~ and clean.

① gene clean (from product of PCR)

② gel clean through column (according introduction of manufacturer) (50ul of PCR product
two tube one is pellet (store in -20°C)
another ~~is~~ has 20ul water. From this, 10ul of
was take for digestion.

B. digestion:

Not I : 26ul water.
3ul Buffer
1ul Not I

1h 37°C

(II, gene clean
II, column clean)

pellet.

↓

BamHI

26ul H₂O
3ul buffer
1ul BamHI

1h 37°C

Ligation.

Water 13ul

Vector 1ul

5x lig buffer 4ul

T4 ligase 2ul

(BamHI and Not I digest)

Control I

Control II

15ul

"

"

"

x

over night 14°C

NotI digestion:

Vector:

10ul	Vector (concent 3.2ug/ul)
3ul	10x buffer
3ul	BSA
4ul	NotI
10ul	H ₂ O
<hr/>	
30ul	37°C 1.5h

target

II₂ and control (MSP-1)

2ul	H ₂ O
3ul	BSA
3ul	10x Buffer
2ul	Enzyme
<hr/>	
30ul	37°C 1.5h

~~Not~~ BamHI digestion

Vector

BamHI	4ul
Buffer	3ul
water	2.3
<hr/>	
30ul	37°C 1.5h

BamHI

Buffer

water

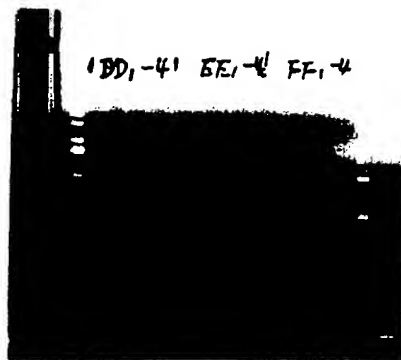
2ul

3ul

25ul

<hr/>	
30ul	37°C 1.5h

Result



FF-4 did not work because first PCR (CC) annealing temp was too high
Need redo CC (first PCR), then FF, FF₄

ligation as before
transformation as before

result. not so much white clones. probably vector
was not properly digested.
clonify further purify vector.

pick up 40 clone grow overnight.

cell PCR: as regular. 10ul cell 94°C 5min.

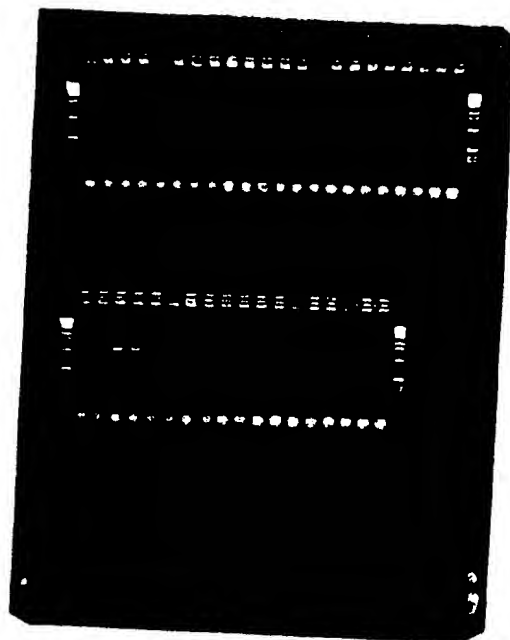
eligo	AL1064	2.5ul
	AB1065	2.5ul
	Buffer	5ul
	dNTP	8ul
	Taq	0.5
	1+20	<u>27.75</u>
		40ul

15 cycle 94°C 45" 50°C 45" 72°C 60"

positive clone

1, 2, 3, 4, 6, 8, 17, 21, 22,
25, 26, 27, 31, 33, 36, 39, 40,

See back



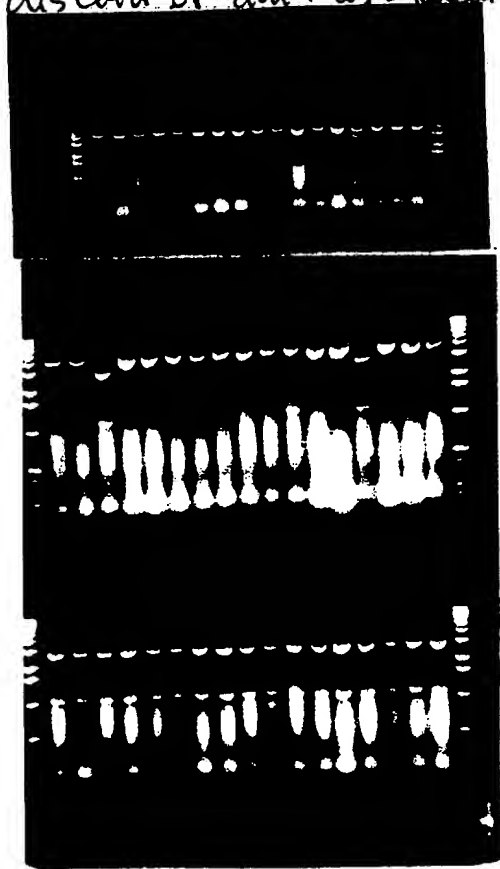
Save cell
Savolas SGV/R⁺ Satellite gene Vaccine
caution - Co sign

digest all positive (17) clones (based on PCR)

Single digestion: BamHI or NotI

double digestion BamHI and NotI.

Result: Clone 3, 26, 33 are not pure clones.
discard or don't use them



2, 6

Plasmid pBacPAK8 and pBacPAK9 (from Sayon
21g/100ul 21g/100ul

Transformation:

10ul plasmide (200ng)

100ul XL-blue cell

procedure as regular.

plating: overnight
growth well

Miniprep of pBacPAK8 and pBacPAK9 -

run undigested and digested plasmid

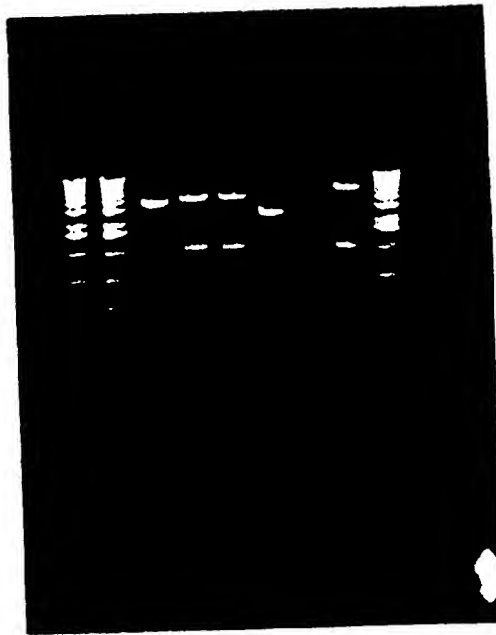


100ng/ μ l $\times 19 =$

1.9ul.

13.2-14

This result confirm that. ~~no~~ orders are
no problem. do confirm (11) (20) (63) clones
are true clones:



I will sequence clone 20.

Methylation:

Clone 63 Vector correct.

Clone 20. Most target correct.

Clone 63 methylation.

Reaction:	3ul	TaqI methylation
	3ul	NEB 4 Buffer
	0.3ul	BSA
	22.2ul	H ₂ O
	1.5ul	Mix SAM

1 hr 65°C

Mix: 50ul NEB 4 Buffer + 450ul H₂O + 1.25ul SAM

0.6ul Nac. (SM)

60ul Ethanol (100%)

Hind II cut

Clone 63 (two piece vector)

Clone 20 (mix of 1000
vector or small)

Run gel Standard

20

63

Standard

(more small) (two big)

15

restriction correlation:

3ul	buffer
8ul	Hind I
21ul	H2O

1.5 hr 37°C

Result:

clone 20

clone 63

|||||

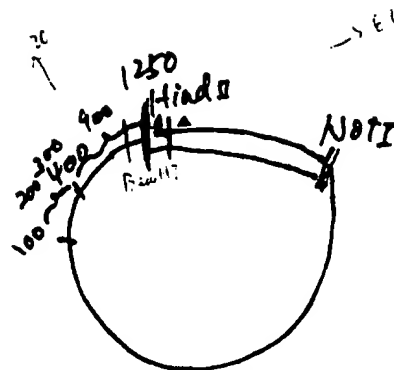
—

Δ

Δ 1Kb
— 0.9Kb

— — 1Kb

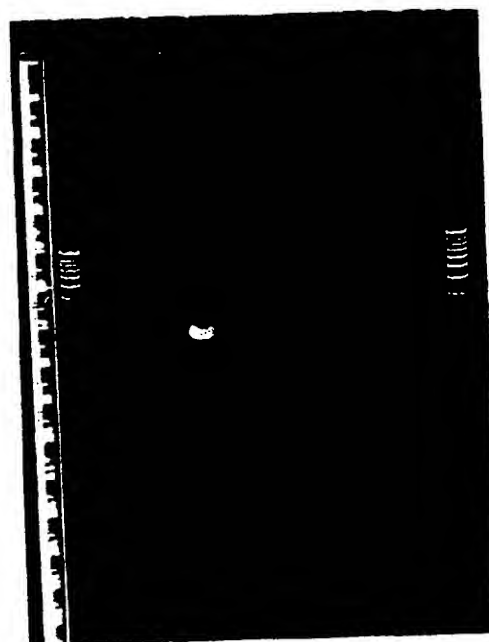
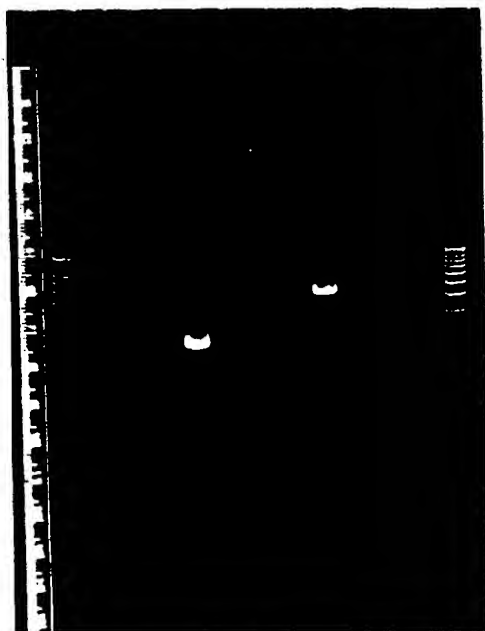
— 0.2 Kb



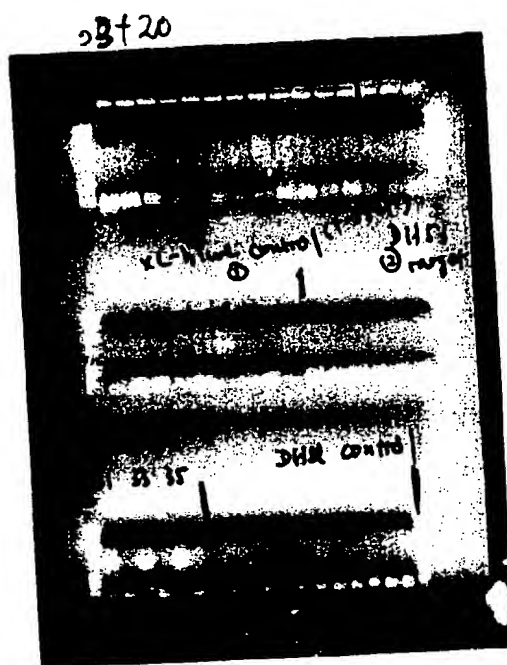
Δ cut fragment

ligation: as routine

100-100



clone 63 + 20 ligation see before
clone PCR primer: AL1097
AL1064



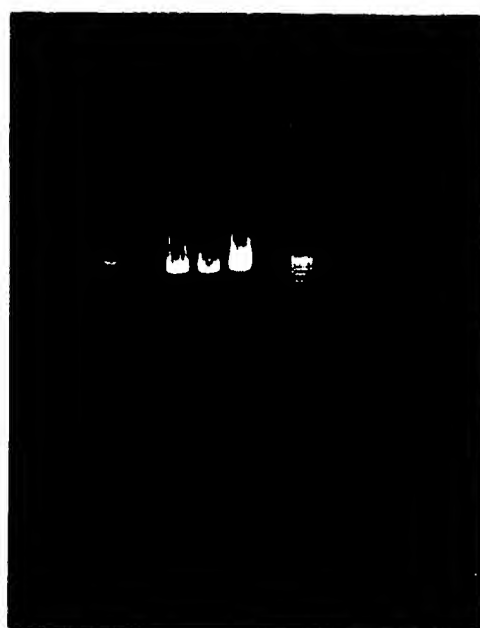
clones 21, 31, 33, 35
are positive

save as name:

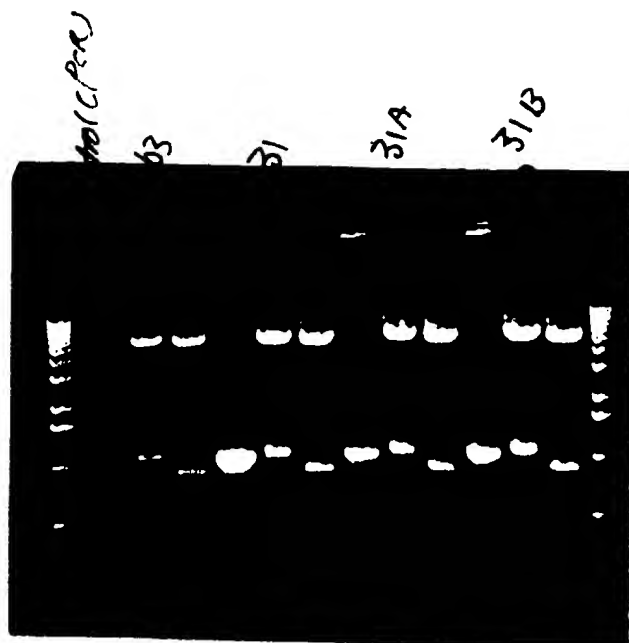
~~38cTS/EL-DT21/63+20~~

hamb
pacB/63+20/number

Bauchtt. digestion: 63+20 (21, 31, 33, 35) 63



confirm primary methylation

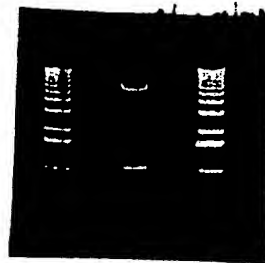


Save clone 31A and 31B
named as pac8/63+20/31A and 31B

Prep for sending product to Hassan.

clone puc8/63+2/31A

BamHI and NotI digested



Send to Hassan